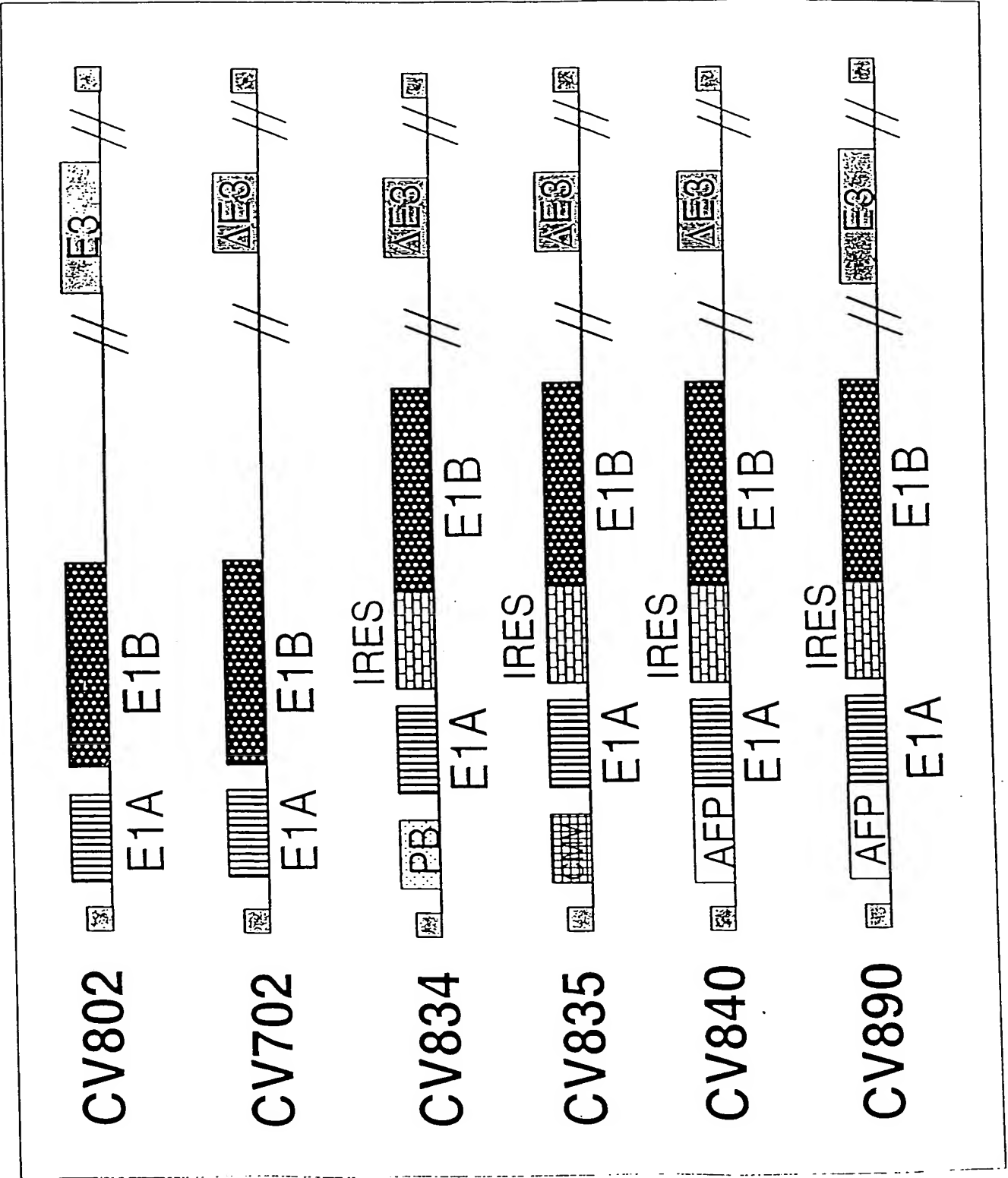


Figure 1

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Figure 2A



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Structure of CV876, CV882 and CV884

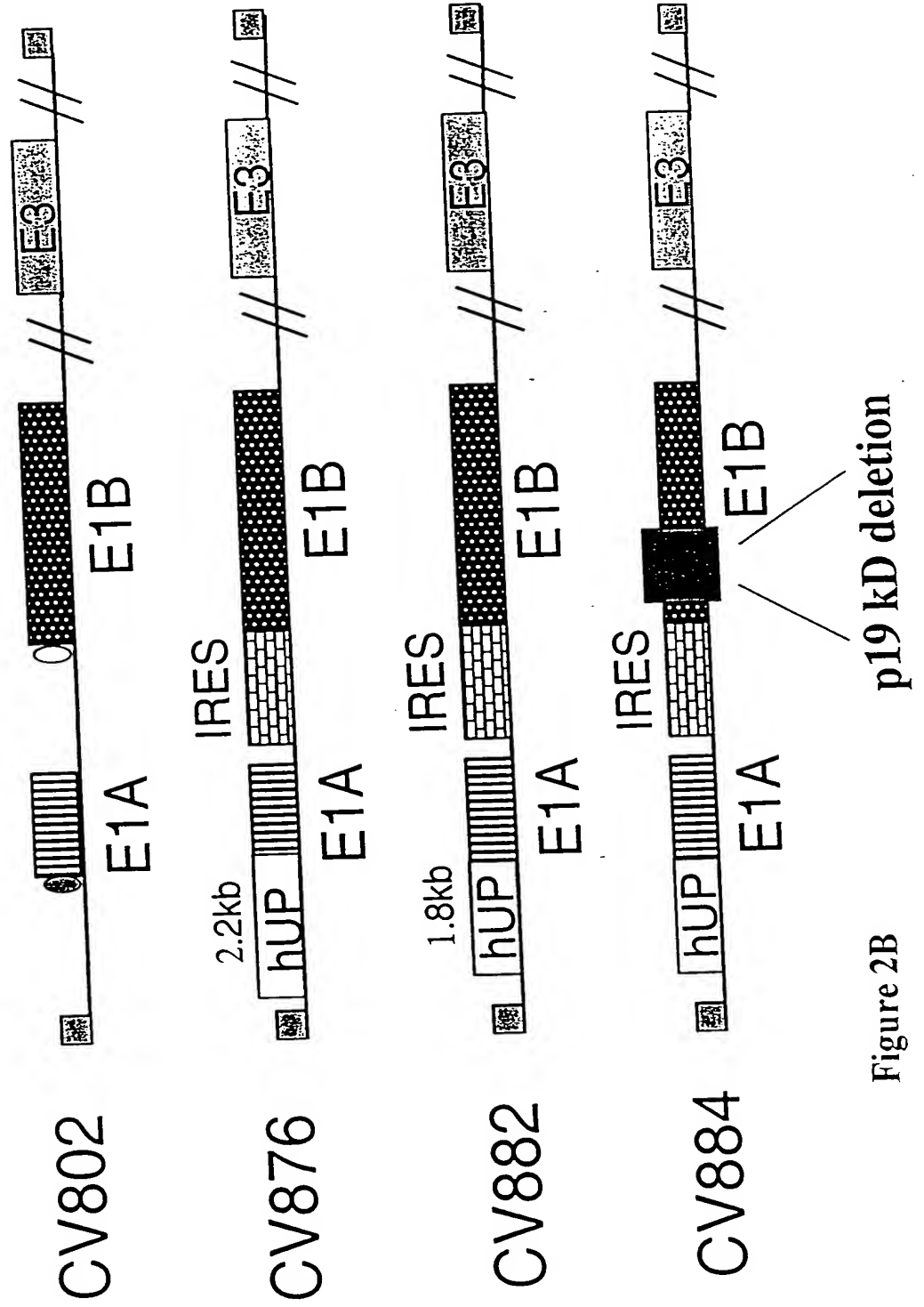


Figure 2B

REPLICATION

293 LNCap HepG2

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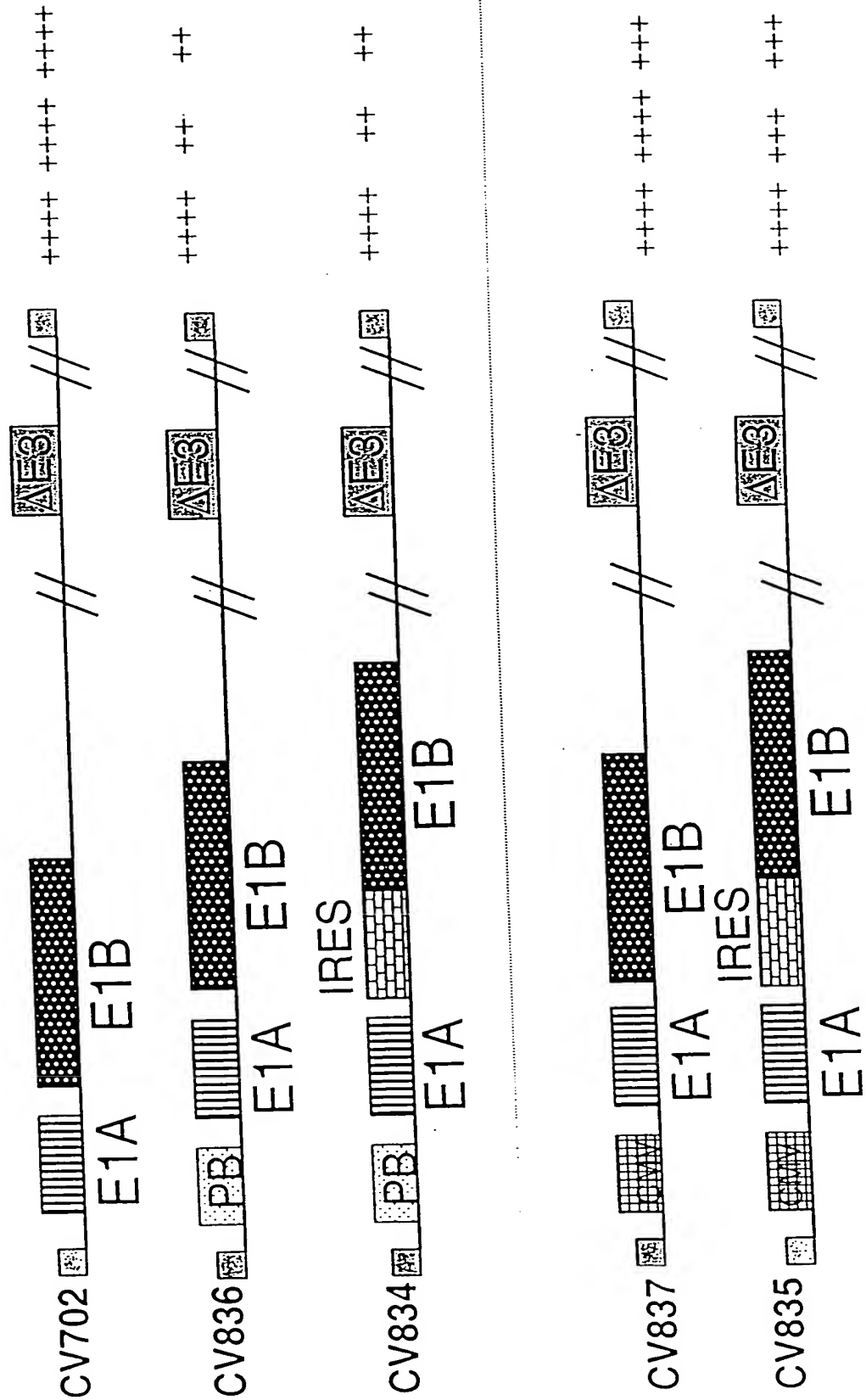
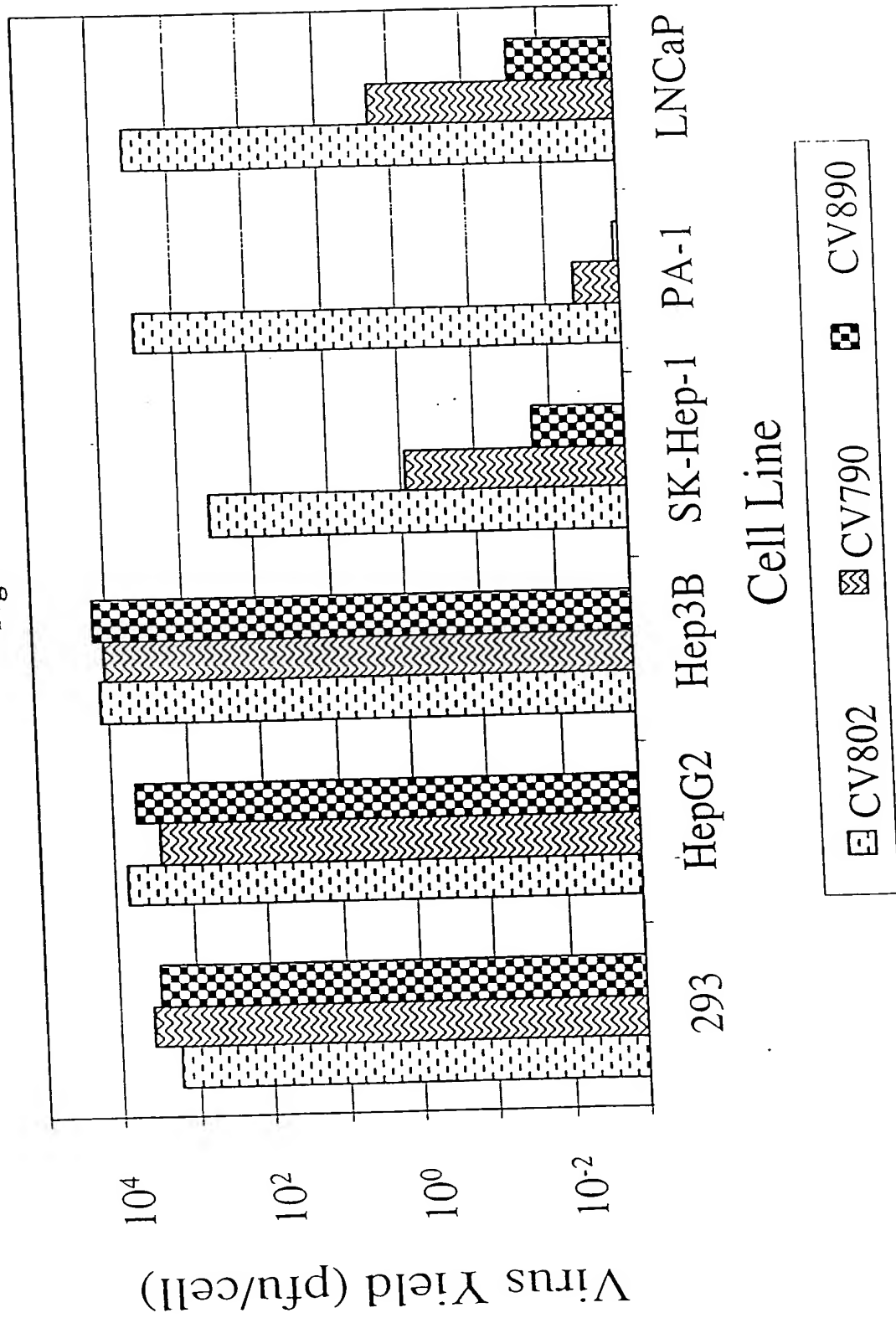


Figure 3

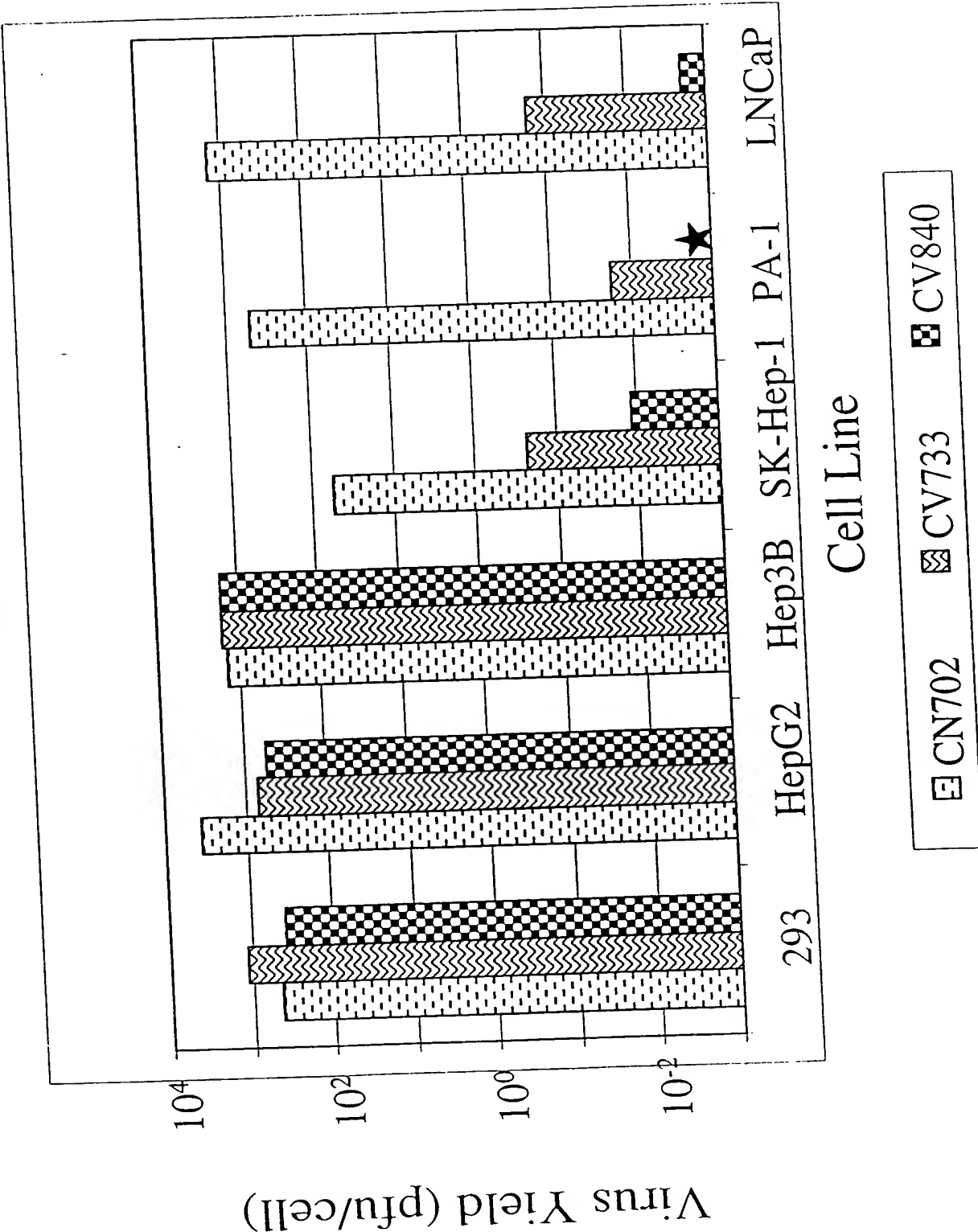
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Figure 4A



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Figure 4B



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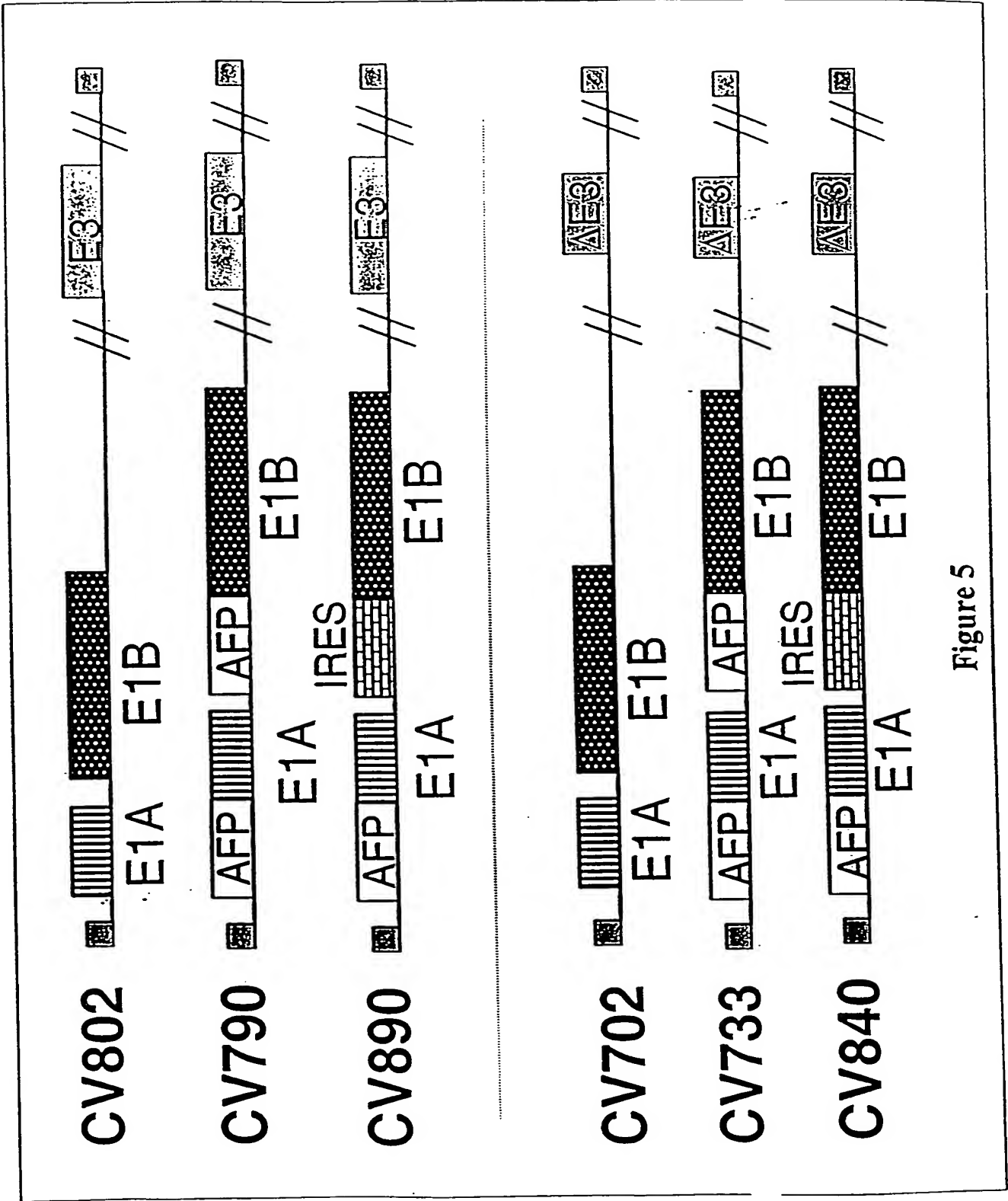
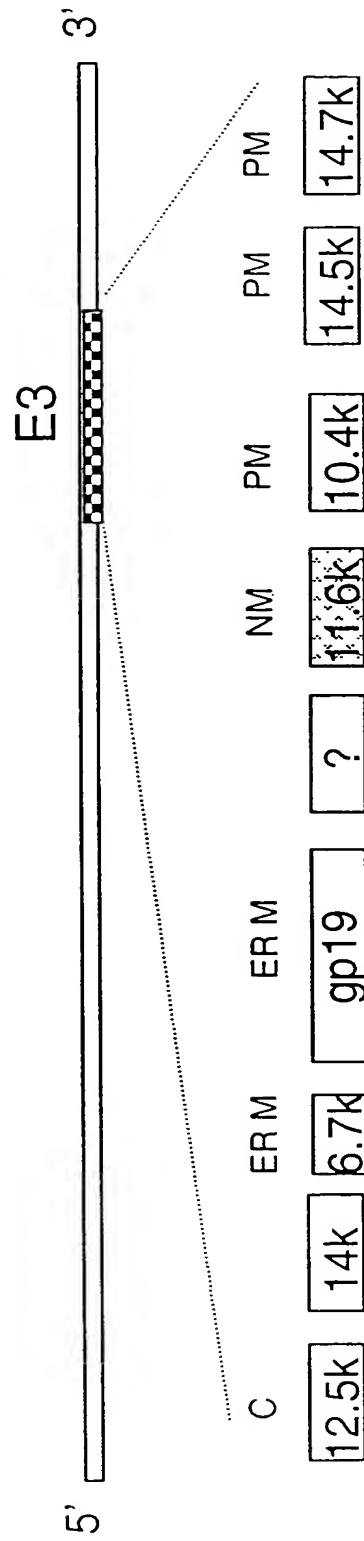


Figure 5

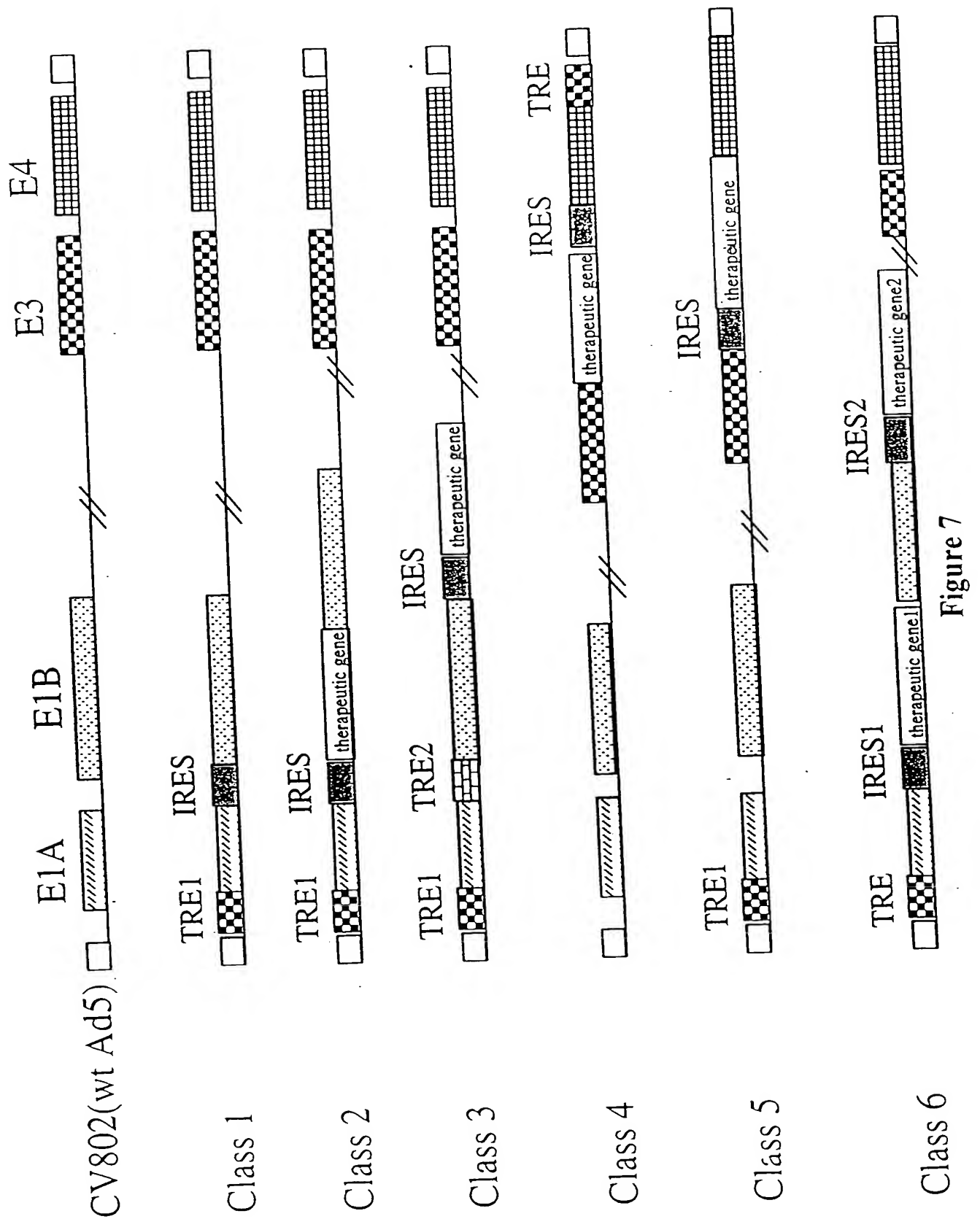
Region E3 of Adenovirus



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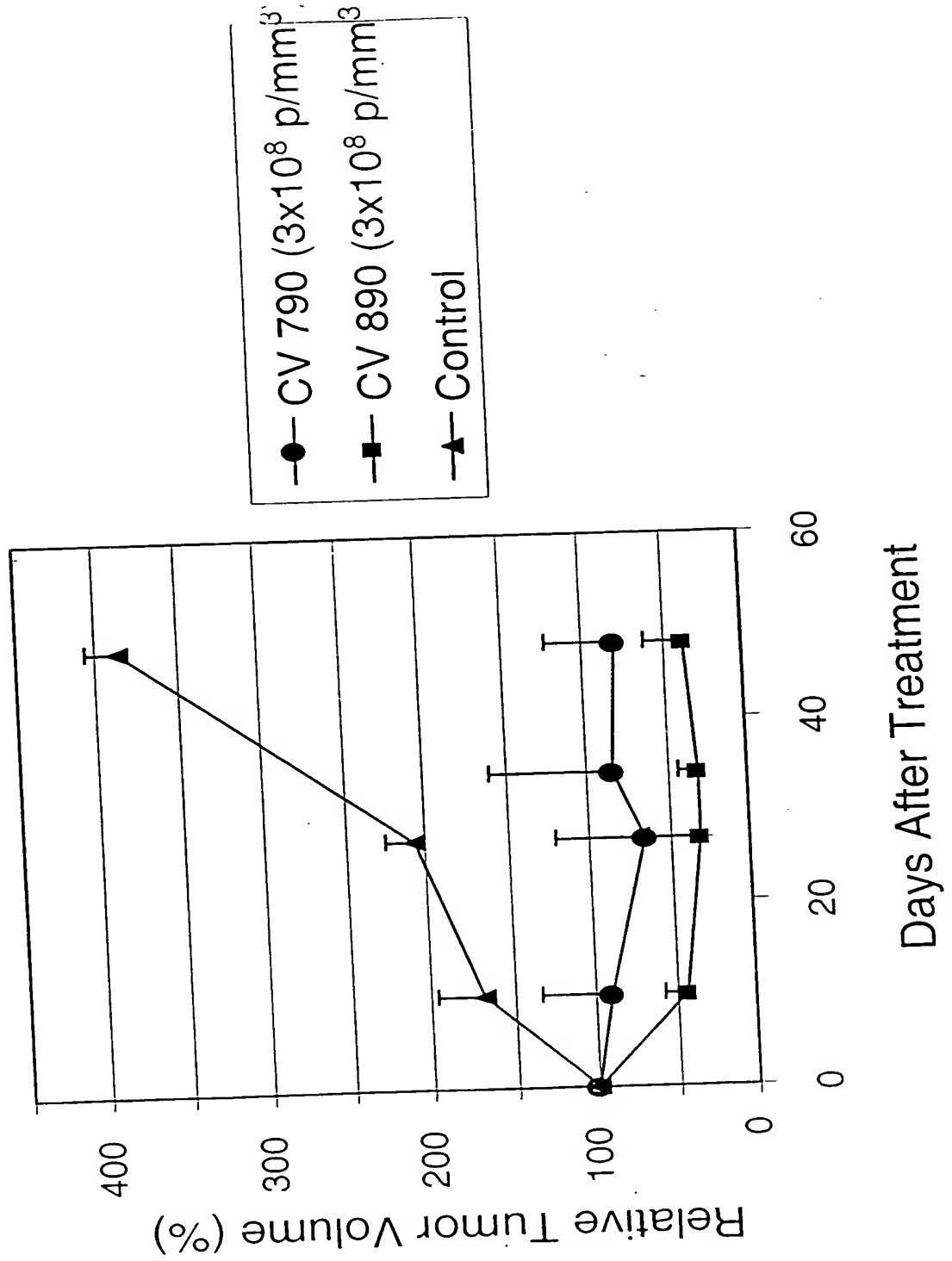
Figure 6

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Figure 8



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FIG. 9

G ATG ACC GGC TCA ACC ATC GCG CCC ACA ACG GAC TAT CGC AAC ACC
 46
 Met. Thr Gly Ser Thr Ile Ala Pro Thr Thr Asp Tyr Arg Asn Thr
 1 5 10 15
 ACT GCT ACC GGA CTA ACA TCT GCC CTA AAT TTA CCC CAA GTT CAT GCC
 94
 Thr Ala Thr Gly Leu Thr Ser Ala Leu Asn Leu Pro Gln Val His Ala
 20 25 30
 TTT GTC AAT GAC TGG GCG AGC TTG GAC ATG TGG TGG TTT TCC ATA GCG
 142
 Phe Val Asn Asp Trp Ala Ser Leu Asp Met Trp Trp Phe Ser Ile Ala
 35 40 45
 CTT ATG TTT GTT TGC CTT ATT ATT ATG TGG CTT ATT TGT TGC CTA AAG
 190
 Leu Met Phe Val Cys Leu Ile Ile Met Trp Leu Ile Cys Cys Leu Lys
 50 55 60
 CGC AGA CGC GCC AGA CCC CCC ATC TAT AGG CCT ATC ATT GTG CTC AAC
 238
 Arg Arg Arg Ala Arg Pro Pro Ile Tyr Arg Pro Ile Ile Val Leu Asn
 65 70 75
 CCA CAC AAT GAA AAA ATT CAT AGA TTG GAC GGT CTG AAA CCA TGT TCT
 286
 Pro His Asn Glu Lys Ile His Arg Leu Asp Gly Leu Lys Pro Cys Ser
 80 85 90 95
 CTT CTT TTA CAG TAT GAT TAA
 307
 Leu Leu Leu Gln Tyr Asp
 100

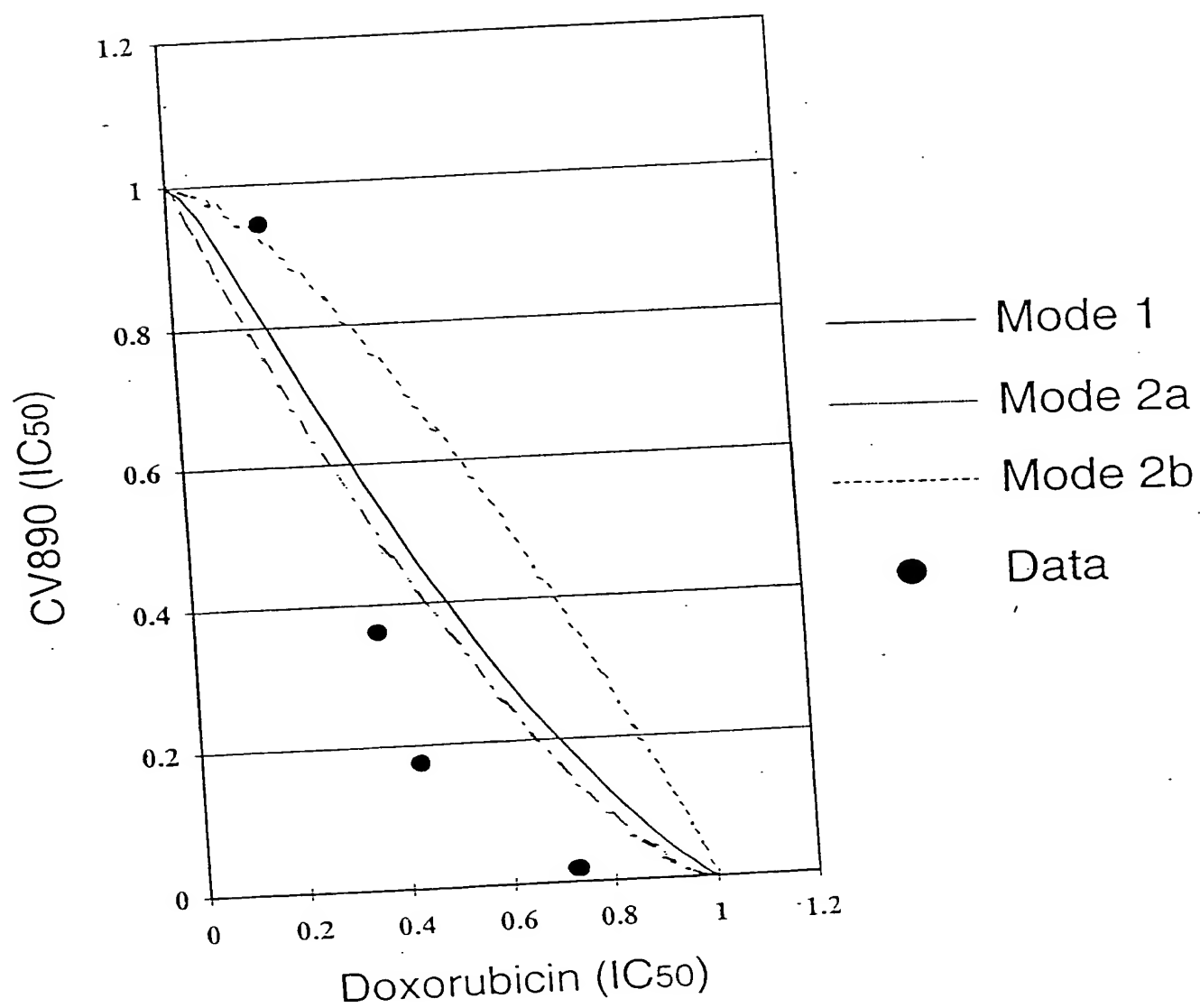


Figure 10

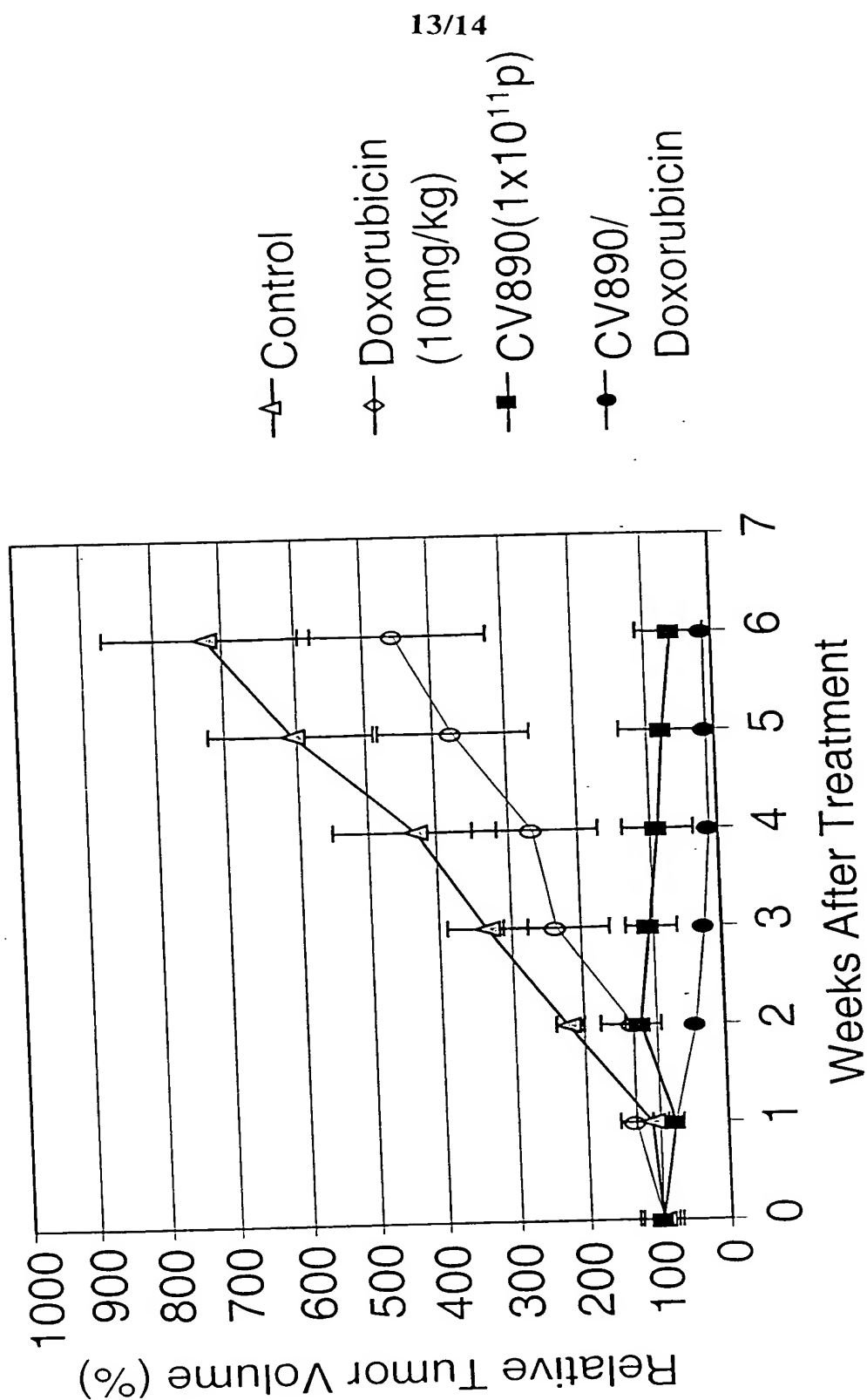


Figure 11

Figure 12

Virus Yield of CV884

